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Terms	Documents
wo adj 9613149	3

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Derwent World Patents Index
 IBM Technical Disclosure Bulletins

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DATE: Tuesday, July 06, 2004 [Printable Copy](#) [Create Case](#)

<u>Set</u> <u>Name</u>	<u>Query</u>	<u>Hit</u> <u>Count</u>	<u>Set</u> <u>Name</u> result set
side by side			
<i>DB=DWPI; PLUR=YES; OP=OR</i>			
<u>L30</u>	wo adj 9613149	3	<u>L30</u>
<i>DB=PGPB,USPT; PLUR=YES; OP=OR</i>			
<u>L29</u>	wo adj 9613149	0	<u>L29</u>
<u>L28</u>	potrykus-Ingo\$.in.	10	<u>L28</u>
<i>DB=USPT; PLUR=YES; OP=OR</i>			
<u>L27</u>	L24 and desaturase and synthase	0	<u>L27</u>
<u>L26</u>	L24 and desaturase and synthase.clm.	0	<u>L26</u>
<u>L25</u>	L24 and desaturase.clm. and synthase.clm.	0	<u>L25</u>
<u>L24</u>	potrykus-Ingo\$.in.	9	<u>L24</u>

<u>L23</u>	5705624.pn.	1	<u>L23</u>
	<i>DB=DWPI; PLUR=YES; OP=OR</i>		
<u>L22</u>	wo adj 9613149	3	<u>L22</u>
<u>L21</u>	wo adj 9413149	1	<u>L21</u>
	<i>DB=USPT; PLUR=YES; OP=OR</i>		
<u>L20</u>	4727219.pn.	1	<u>L20</u>
<u>L19</u>	4727219	433	<u>L19</u>
	<i>DB=EPAB; PLUR=YES; OP=OR</i>		
<u>L18</u>	WO-9806862-A1.did.	1	<u>L18</u>
<u>L17</u>	WO-9806862-A1.did.	1	<u>L17</u>
<u>L16</u>	WO-9806862-A1.did.	1	<u>L16</u>
	<i>DB=DWPI; PLUR=YES; OP=OR</i>		
<u>L15</u>	wo adj 9806862	1	<u>L15</u>
<u>L14</u>	wo adj 98/06862	0	<u>L14</u>
<u>L13</u>	L10 and synthase	1	<u>L13</u>
<u>L12</u>	L10 and desaturase and synthase	0	<u>L12</u>
<u>L11</u>	L10 and desaturase and synthase and phytoene	0	<u>L11</u>
<u>L10</u>	wo adj 9907867	1	<u>L10</u>
<u>L9</u>	wo adj 99/07867	0	<u>L9</u>
	<i>DB=USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR</i>		
<u>L8</u>	phytoene and desaturase and synthase and carotene and carotenoid and rice	25	<u>L8</u>
<u>L7</u>	phytoene and desaturase and synthase and plant and carotene and carotenoid and rice	25	<u>L7</u>
<u>L6</u>	phytoene and desaturase and synthase and plant and carotene	75	<u>L6</u>
<u>L5</u>	phytoene.clm and desaturase and synthase and plant	0	<u>L5</u>
<u>L4</u>	phytoene and desaturase and synthase and plant	112	<u>L4</u>
<u>L3</u>	phytoene and desaturase and synthase.clm and plant	0	<u>L3</u>
<u>L2</u>	phytoene and desaturase.clm and synthase.clm and plant	0	<u>L2</u>
<u>L1</u>	phytoene.clm and desaturase.clm and synthase.clm and plant	0	<u>L1</u>

END OF SEARCH HISTORY

COST IN U.S. DOLLARS	SINCE FILE	TOTAL
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FULL ESTIMATED COST	0.21	0.21

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=> s phytoene and desaturase and synthase and carotenoid
L1 191 PHYTOENE AND DESATURASE AND SYNTHASE AND CAROTENOID

=> duplicate remove l1
DUPLICATE PREFERENCE IS 'AGRICOLA, BIOSIS, EMBASE, CAPLUS'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n

PROCESSING COMPLETED FOR L1
L2 104 DUPLICATE REMOVE L1 (87 DUPLICATES REMOVED)

=> s l2 phytoene(w)synthase
MISSING OPERATOR L2 PHYTOENE
The search profile that was entered contains terms or
nested terms that are not separated by a logical operator.

=> s l2 and phytoene(w)synthase
L3 100 L2 AND PHYTOENE(W) SYNTHASE

=> s l3 and phytoene(w)desaturase
L4 93 L3 AND PHYTOENE(W) DESATURASE

=> s l3 and plant and transform?
L5 14 L3 AND PLANT AND TRANSFORM?

=> d l5 1-15 ibib ab

L5 ANSWER 1 OF 14 AGRICOLA Compiled and distributed by the National
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of America. It contains copyrighted materials. All rights reserved.
(2004) on STN

ACCESSION NUMBER:	2003:27813 AGRICOLA
DOCUMENT NUMBER:	IND23315255
TITLE:	An evaluation of factors affecting the efficiency of Agrobacterium-mediated ***transformation*** of Citrus paradisi (Macf.) and production of transgenic ***plants*** containing ***carotenoid*** biosynthetic genes.
AUTHOR(S):	Costa, M.G.C.; Otoni, W.C.; Moore, G.A.
SOURCE:	Plant cell reports, Nov 2002. Vol. 21, No. 4. p. 365-373

Publisher: Berlin : Springer-Verlag.

CODEN: PCRPD8; ISSN: 0721-7714

NOTE:

Includes references

PUB. COUNTRY:

Germany

DOCUMENT TYPE:

Article

FILE SEGMENT:

Non-U.S. Imprint other than FAO

LANGUAGE:

English

AB An improved protocol for Agrobacterium-mediated ***transformation*** of Duncan grapefruit (Citrus paradisi Macf.) epicotyl explants was developed by examining the effects of six different factors on the efficiency of ***transformation*** and combining the best treatment for each factor. The preculturing of explants and the composition of the cocultivation medium were the factors that most influenced ***transformation*** efficiency. The optimized protocol was successfully employed in the production of transgenic grapefruit ***plants*** containing the ***carotenoid*** biosynthetic genes ***phytoene*** ***synthase***, ***phytoene*** ***desaturase***, or lycopene-beta-cyclase under constitutive expression. With an eventual goal of metabolically engineering grapefruit with multiple genes, hygromycin as a selectable marker and BIBAC as a ***transformation*** vector for large pieces of DNA were also tested.

L5 ANSWER 2 OF 14 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2004) on STN

ACCESSION NUMBER:

2002:47449 AGRICOLA

DOCUMENT NUMBER:

IND23281522

TITLE:

Functional analysis of the early steps of ***carotenoid*** biosynthesis in tobacco.

AUTHOR(S):

Busch, M.; Seuter, A.; Hain, R.

AVAILABILITY:

DNAL (450 P692)

SOURCE:

Plant physiology, Feb 2002. Vol. 128, No. 2. p. 439-453

Publisher: Rockville, MD : American Society of Plant Physiologists, 1926-

CODEN: PLPHAY; ISSN: 0032-0889

NOTE:

Includes references

PUB. COUNTRY:

Maryland; United States

DOCUMENT TYPE:

Article; Conference

FILE SEGMENT:

U.S. Imprints not USDA, Experiment or Extension

LANGUAGE:

English

AB ***Carotenoids*** contribute to energy transduction in the light harvesting complexes and serve in protection from excess light fluence. Because of the importance of ***carotenoids***, the genes encoding enzymes of ***carotenoid*** biosynthesis in higher ***plants*** are potential targets for herbicides. To obtain further insight into tobacco ***carotenoid*** biosynthesis and to investigate and prioritize potential herbicide targets in the pathway, the effects of changed ***phytoene*** ***synthase*** (PSY) and ***phytoene*** ***desaturase*** (PDS) gene expression were studied in transgenic tobacco (Nicotiana tabacum Petit Havana SR1) ***plants***. Genes for both enzymes were cloned from tobacco, and surprisingly two functional PSY genes were found. Transgenic tobacco ***plants*** constitutively expressing these genes in both sense and antisense orientations were examined regarding phenotype, ***carotenoid*** content and transcript

levels of carotene biosynthesis genes. Overexpression of either psy gene resulted in severe phenotypic effects including dwarfism, altered leaf morphology, and pigmentation. A correlation among phenotype, transcript level, and metabolic profile was demonstrated by comparison of hemizygous and homozygous ***plants*** from the same ***transformation*** event. Antisense expression of PSY and PDS also caused lethal phenotypes. Transcript levels of other carotene biosynthesis genes remained unaltered in the transgenic mutant. ***Phytoene*** accumulated in ***plants*** expressing antisense RNA to pds. However, elevated levels of ***phytoene*** were detected suggesting an increase in metabolic flux into this pathway.

L5 ANSWER 3 OF 14 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
 ACCESSION NUMBER: 2004:107975 BIOSIS
 DOCUMENT NUMBER: PREV200400110881
 TITLE: Metabolic engineering of the astaxanthin-biosynthetic pathway of Xanthophyllomyces dendrorhous.
 AUTHOR(S): Visser, Hans [Reprint Author]; van Ooyen, Albert J. J.; Verdoes, Jan C.
 CORPORATE SOURCE: Section of Fungal Genomics, Wageningen University, Dreijenlaan 2, 6703 HA, Wageningen, Netherlands
 hans.visser@wur.nl
 SOURCE: FEMS Yeast Research, (December 2003) Vol. 4, No. 3, pp. 221-231. print.
 ISSN: 1567-1356 (ISSN print).
 DOCUMENT TYPE: Article
 General Review; (Literature Review)
 LANGUAGE: English
 ENTRY DATE: Entered STN: 25 Feb 2004
 Last Updated on STN: 25 Feb 2004

AB This review describes the different approaches that have been used to manipulate and improve ***carotenoid*** production in Xanthophyllomyces dendrorhous. The red yeast X. dendrorhous (formerly known as Phaffia rhodozyma) is one of the microbiological production systems for natural astaxanthin. Astaxanthin is applied in food and feed industry and can be used as a nutraceutical because of its strong antioxidant properties. However, the production levels of astaxanthin in wild-type isolates are rather low. To increase the astaxanthin content in X. dendrorhous, cultivation protocols have been optimized and astaxanthin-hyperproducing mutants have been obtained by screening of classically mutagenized X. dendrorhous strains. The knowledge about the regulation of carotenogenesis in X. dendrorhous is still limited in comparison to that in other carotenogenic fungi. The X. dendrorhous carotenogenic genes have been cloned and a X. dendrorhous ***transformation*** system has been developed. These tools allowed

the

directed genetic modification of the astaxanthin pathway in X. dendrorhous. The crtYB gene, encoding the bifunctional enzyme ***phytoene*** ***synthase*** /lycopene cyclase, was inactivated by insertion of a vector by single and double cross-over events, indicating that it is possible to generate specific ***carotenoid*** -biosynthetic mutants. Additionally, overexpression of crtYB resulted in the accumulation of beta-carotene and echinone, which indicates that the oxygenation reactions are rate-limiting in these recombinant strains. Furthermore, overexpression of the ***phytoene*** ***desaturase*** -encoding gene (crt1) showed an increase in monocyclic ***carotenoids*** such as torulene and HDCO (3-hydroxy-3',4'-didehydro-beta-psi-carotene-4-

one) and a decrease in bicyclic ***carotenoids*** such as echinone, beta-carotene and astaxanthin.

L5 ANSWER 4 OF 14 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
ACCESSION NUMBER: 2003:397089 BIOSIS
DOCUMENT NUMBER: PREV200300397089
TITLE: Metabolic engineering of the ***carotenoid***
biosynthetic pathway in the yeast *Xanthophyllomyces dendrorhous* (*Phaffia rhodozyma*).
AUTHOR(S): Verdoes, Jan C.; Sandmann, Gerhard; Visser, Hans [Reprint
Author]; Diaz, Maria; van Mossel, Minca; van Ooyen, Albert
J. J.
CORPORATE SOURCE: Laboratory of Microbiology, Section of Fungal Genomics,
Department of Agrotechnology and Food Sciences, Wageningen
University, Dreijenlaan 2, 6703 HA, Wageningen, Netherlands
hans.visser@wur.nl
SOURCE: Applied and Environmental Microbiology, (July 2003) Vol.
69, No. 7, pp. 3728-3738. print.
ISSN: 0099-2240 (ISSN print).
DOCUMENT TYPE: Article
LANGUAGE: English
ENTRY DATE: Entered STN: 27 Aug 2003
Last Updated on STN: 27 Aug 2003

AB The crtYB locus was used as an integrative platform for the construction of specific ***carotenoid*** biosynthetic mutants in the astaxanthin-producing yeast *Xanthophyllomyces dendrorhous*. The crtYB gene of *X. dendrorhous*, encoding a chimeric ***carotenoid*** biosynthetic enzyme, could be inactivated by both single and double crossover events, resulting in non- ***carotenoid*** -producing ***transformants***. In addition, the crtYB gene, linked to either its homologous or a glyceraldehyde-3-phosphate dehydrogenase promoter, was overexpressed in the wild type and a beta-carotene-accumulating mutant of *X. dendrorhous*. In several ***transformants*** containing multiple copies of the crtYB gene, the total ***carotenoid*** content was higher than in the control strain. This increase was mainly due to an increase of the beta-carotene and echinone content, whereas the total content of astaxanthin was unaffected or even lower. Overexpression of the ***phytoene*** ***synthase*** -encoding gene (crtI) had a large impact on the ratio between mono- and bicyclic ***carotenoids***. Furthermore, we showed that in metabolically engineered *X. dendrorhous* strains, the competition between the enzymes ***phytoene*** ***desaturase*** and lycopene cyclase for lycopene governs the metabolic flux either via beta-carotene to astaxanthin or via 3,4-didehydrolycopene to 3-hydroxy-3'-4'-didehydro-beta-psi-caroten-4-one (HDCO). The monocyclic ***carotenoid*** torulene and HDCO, normally produced as minority ***carotenoids***, were the main ***carotenoids*** produced in these strains.

L5 ANSWER 5 OF 14 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
ACCESSION NUMBER: 2000:335140 BIOSIS
DOCUMENT NUMBER: PREV200000335140
TITLE: Elevation of the provitamin A content of transgenic tomato
plants.
AUTHOR(S): Romer, Susanne; Fraser, Paul D.; Kiano, Joy W.; Shipton,
Cathie A.; Misawa, Norihiko; Schuch, Wolfgang; Bramley,

Peter M. [Reprint author]
 CORPORATE SOURCE: School of Biological Sciences, Royal Holloway, University
 of London, Egham, Surrey, TW20 OEX, UK
 SOURCE: Nature Biotechnology, (June, 2000) Vol. 18, No. 6, pp.
 666-669. print.
 ISSN: 1087-0156.
 DOCUMENT TYPE: Article
 LANGUAGE: English
 ENTRY DATE: Entered STN: 10 Aug 2000
 Last Updated on STN: 7 Jan 2002

AB Tomato products are the principal dietary sources of lycopene and major
 source of beta-carotene, both of which have been shown to benefit human
 health. To enhance the ***carotenoid*** content and profile of tomato
 fruit, we have produced transgenic lines containing a bacterial
 carotenoid gene (crtI) encoding the enzyme ***phytoene***
 desaturase, which converts ***phytoene*** into lycopene.
 Expression of this gene in transgenic tomatoes did not elevate total
 carotenoid levels. However, the beta-carotene content increased
 about threefold, up to 45% of the total ***carotenoid*** content.
 Endogenous ***carotenoid*** genes were concurrently upregulated,
 except for ***phytoene*** ***synthase***, which was repressed.
 The alteration in ***carotenoid*** content of these ***plants***
 did not affect growth and development. Levels of noncarotenoid
 isoprenoids were unchanged in the ***transformants***. The phenotype
 has been found to be stable and reproducible over at least four
 generations.

L5 ANSWER 6 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 2003:922684 CAPLUS
 DOCUMENT NUMBER: 140:1581
 TITLE: Regulation of genes involved in ***carotenoid***
 and tocopherol biosynthesis pathway in transgenic
 plants for producing ***carotenoid***
 compounds, tocopherol compounds, and specialty oils in
 plant seeds
 INVENTOR(S): Shewmaker, Christine K.; Bhat, B. Ganesh;
 Venkatramesh, Mylavaraapu; Rangwala, Shaikat H.;
 Kishore, Ganesh M.; Boddupalli, Sekhar S.
 PATENT ASSIGNEE(S): Calgene LLC, USA
 SOURCE: U.S., 57 pp.
 CODEN: USXXAM
 DOCUMENT TYPE: Patent
 LANGUAGE: English
 FAMILY ACC. NUM. COUNT: 1
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 6653530	B1	20031125	US 1998-23587	19980213
PRIORITY APPLN. INFO.:			US 1998-23587	19980213
AB Methods are provided for producing ***plants*** and seeds having altered ***carotenoid***, fatty acid and tocopherol compns. The methods find particular use in increasing the ***carotenoid*** and tocopherol levels in oilseed ***plants***, and in providing desirable high oleic acid seed oils. Specifically, chimeric genes encoding E. uredovora ***phytoene*** ***synthase*** (crtB), or ***phytoene*** ***desaturase*** (crtI), or GGPP ***synthase***				

(crtE) in fusion with plastid transit peptide of pea Rubisco small subunit (rbcS) under the control of seed-preferred napin gene promoter, and napin or nos termination region, are constructed to make transgenic Brassica napus. Binary constructs expressing both crtB and crtI genes, crtB and antisense epsilon cyclase or beta cyclase genes are also used to
 transform Brassica napus. Also demonstrated are increased
 carotenoid prodn., particularly increased ratio of
 .alpha.-carotene and .beta.-carotene to ***phytoene***, and increased levels of oleic acid and decreased levels of linoleic and/or linolenic acid in seeds of transgenic ***plants***.

REFERENCE COUNT: 104 THERE ARE 104 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE REFORMAT

L5 ANSWER 7 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 2003:920076 CAPLUS

DOCUMENT NUMBER: 140:250219

TITLE: Coordinate expression of multiple bacterial
 carotenoid genes in canola leading to altered
 carotenoid production

AUTHOR(S): Ravanello, Monica P.; Ke, Dangyang; Alvarez, Julie; Huang, Bihua; Shewmaker, Christine K.

CORPORATE SOURCE: Calgene Campus, Monsanto Company, Davis, CA, 95616, USA

SOURCE: Metabolic Engineering (2003), 5(4), 255-263
 CODEN: MEENFM; ISSN: 1096-7176

PUBLISHER: Elsevier Science

DOCUMENT TYPE: Journal

LANGUAGE: English

AB ***Carotenoids*** have drawn much attention recently because of their potentially pos. benefits to human health as well as their utility in both food and animal feed. Previous work in canola (Brassica napus) seed over-expressing the bacterial ***phytoene*** ***synthase*** gene (crtB) demonstrated a change in ***carotenoid*** content, such that the total levels of ***carotenoids***, including ***phytoene*** and downstream metabolites like .beta.-carotene, were elevated 50-fold, with the ratio of .beta.- to .alpha.-carotene being 2:1. This result raised the possibility that the compn. of metabolites in this pathway could be modified further in conjunction with the increased flux obtained with crtB. Here we report on the expression of addnl. bacterial genes for the enzymes geranylgeranyl diphosphate ***synthase*** (crtE), ***phytoene*** ***desaturase*** (crtI) and lycopene cyclase (crtY and the ***plant*** B. napus lycopene .beta.-cyclase) engineered in conjunction with ***phytoene*** ***synthase*** (crtB) in transgenic canola seed. Anal. of the ***carotenoid*** levels by HPLC revealed a 90% decrease in ***phytoene*** levels for the double construct expressing crtB in conjunction with crtI. The transgenic seed from all the double constructs, including the one expressing the bacterial crtB and the ***plant*** lycopene .beta.-cyclase showed an increase in the levels of total ***carotenoid*** similar to that previously obsd. by expressing crtB alone but minimal effects were obsd. with respect to the ratio of .beta.- to .alpha.-carotene compared to the original construct. However, the .beta.- to .alpha.-carotene ratio was increased from 2:1 to 3:1 when a triple construct consisting of the bacterial
 phytoene ***synthase***, ***phytoene***
 desaturase

and lycopene cyclase genes were expressed together. This result suggests

that the bacterial genes may form an aggregate complex that allows in vivo activity of all three proteins through substrate channeling. This finding should allow further manipulation of the ***carotenoid*** biosynthetic pathway for downstream products with enhanced agronomic, animal feed and human nutritional values.

REFERENCE COUNT: 28 THERE ARE 28 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L5 ANSWER 8 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 2003:909740 CAPLUS

DOCUMENT NUMBER: 140:214068

TITLE: Bioengineered 'golden' indica rice cultivars with .beta.-carotene metabolism in the endosperm with hygromycin and mannose selection systems

AUTHOR(S): Datta, Karabi; Baisakh, Niranjan; Oliva, Norman; Torrizo, Lina; Abrigo, Editha; Tan, Jing; Rai, Mayank; Rehana, Sayda; Al-Babili, Salim; Beyer, Peter; Potrykus, Ingo; Datta, Swapan K.

CORPORATE SOURCE: Plant Breeding, Genetics, and Biochemistry Division, International Rice Research Institute, Metro Manila, Philippines

SOURCE: Plant Biotechnology Journal (2003), 1(2), 81-90
CODEN: PBJLAE; ISSN: 1467-7644

PUBLISHER: Blackwell Publishing Ltd.

DOCUMENT TYPE: Journal

LANGUAGE: English

AB Vitamin-A deficiency (VAD) is a major malnutrition problem in South Asia, where indica rice is the staple food. Indica-type rice varieties feed more than 2 billion people. Hence, the authors introduced a combination of transgenes using the biolistic system of ***transformation*** enabling biosynthesis of provitamin A in the endosperm of several indica rice cultivars adapted to diverse ecosystems of different countries. The rice seed-specific glutelin promoter (Gt-1 P) was used to drive the expression of ***phytoene*** ***synthase*** (psy), while lycopene .beta.-cyclase (lcy) and ***phytoene*** ***desaturase*** (crtl), fused to the transit peptide sequence of the pea-Rubisco small subunit, were driven by the constitutive cauliflower mosaic virus promoter (CaMV35S P). Transgenic ***plants*** were recovered through selection with either CaMV35S P driven hph (hygromycin phosphotransferase) gene or cestrum yellow leaf curling virus promoter (CMP) driven pmi (phosphomannose isomerase) gene. Mol. and biochem. analyses demonstrated stable integration and expression of the transgenes. The yellow color of the polished rice grain evidenced the ***carotenoid*** accumulation in the endosperm. The color intensity correlated with the estd.

carotenoid content by spectrophotometric and HPLC anal.

Carotenoid level in cooked polished seeds was comparable (with minor loss of xanthophylls) to that in non-cooked seeds of the same transgenic line. The variable segregation pattern in T1 selfing generation indicated single to multiple loci insertion of the transgenes in the genome. This is the first report of using nonantibiotic pmi driven by a novel promoter in generating transgenic indica rice for possible future use in human nutrition.

REFERENCE COUNT: 43 THERE ARE 43 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L5 ANSWER 9 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 2003:154580 CAPLUS

DOCUMENT NUMBER: 138:199995
 TITLE: Pantoea stewartii genes encoding enzymes involved in
 carotenoid compound conversion from
 phytoene and use thereof
 INVENTOR(S): Brzostowicz, Patricia C.; Cheng, Qiong; Picataggio,
 Stephen K.; Rouviere, Pierre E.
 PATENT ASSIGNEE(S): E. I. Du Pont de Nemours & Co., USA
 SOURCE: PCT Int. Appl., 68 pp.
 CODEN: PIXXD2
 DOCUMENT TYPE: Patent
 LANGUAGE: English
 FAMILY ACC. NUM. COUNT: 1
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2003016503	A2	20030227	WO 2002-US26647	20020815

W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
 CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
 GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR,
 LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH,
 PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ,
 UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD,
 RU, TJ, TM

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, BG,
 CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL,
 PT, SE, SK, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR,
 NE, SN, TD, TG

US 2003148319	A1	20030807	US 2002-218118	20020813
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PRIORITY APPLN. INFO.: US 2001-312646P P 20010815

AB Genes have been isolated from Pantoea stewartii encoding geranylgeranyl
 pyrophosphate (GGPP) ***synthase*** (crtE), ***phytoene***
 synthase (crtB), ***phytoene*** ***desaturase*** (crtl),
 lycopene cyclase (crtY), .beta.-carotene hydroxylase (crtZ), and
 zeaxanthin glucosyl transferase (crtX) activity. The genes and their
 products are useful for the conversion of ***phytoene*** to the
 carotenoids. Vectors contg. those DNA segments, host cells
 contg.

the vectors and methods for producing those enzymes and .beta.-carotene by
 recombinant DNA technol. in ***transformed*** host organisms are
 disclosed.

L5 ANSWER 10 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 2002:172119 CAPLUS

DOCUMENT NUMBER: 136:231339

TITLE: ***Carotenoid*** production from a single carbon
 substrate

INVENTOR(S): Brzostowicz, Patricia C.; Cheng, Qiong; Dicosimo,
 Deana J.; Koffas, Mattheos; Miller, Edward S.; Odom,
 J. Martin; Picataggio, Stephen K.; Rouviere, Pierre E.

PATENT ASSIGNEE(S): E. I. Du Pont de Nemours & Co., USA

SOURCE: PCT Int. Appl., 156 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 4

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
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WO 2002018617	A2	20020307	WO 2001-US27420	20010904
WO 2002018617	A3	20030522		
W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG				
US 2002142408	A1	20021003	US 2001-938956	20010824
US 2003003528	A1	20030102	US 2001-941947	20010829
AU 2001088699	A5	20020313	AU 2001-88699	20010904
EP 1328639	A2	20030723	EP 2001-968453	20010904
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL, TR				
NO 2003000343	A	20030403	NO 2003-343	20030123
US 2004077068	A1	20040422	US 2003-363567	20030904
US 2004063143	A1	20040401	US 2003-700003	20031103
PRIORITY APPLN. INFO.:			US 2000-229858P	P 20000901
			US 2000-229907P	P 20000901
			US 2001-934903	A3 20010822
			WO 2001-US27420	W 20010904
AB A method for the prodn. of ***carotenoid*** compds. is disclosed. The method relies on the use of microorganisms which metabolize single carbon substrates for the prodn. of ***carotenoid*** compds. in high yields. Thus Methylomonas strain 16a was genetically enhanced to produce .beta.-carotene and zeaxanthin from methane.				
L5 ANSWER 11 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN				
ACCESSION NUMBER:		2001:453094 CAPLUS		
DOCUMENT NUMBER:		135:72153		
TITLE:		Moss genes from Physcomitrella patens encoding proteins involved in the synthesis of tocopherols and ***carotenoids***		
INVENTOR(S):		Lerchl, Jens; Renz, Andreas; Ehrhardt, Thomas; Reindl, Andreas; Cirpus, Petra; Bischoff, Friedrich; Frank, Markus; Freund, Annette; Duwenig, Elke; Schmidt, Ralf-Michael; Reski, Ralf; Badur, Ralf		
PATENT ASSIGNEE(S):		BASF Plant Science G.m.b.H., Germany		
SOURCE:		PCT Int. Appl., 123 pp. CODEN: PIXXD2		
DOCUMENT TYPE:		Patent		
LANGUAGE:		English		
FAMILY ACC. NUM. COUNT:		1		
PATENT INFORMATION:				

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
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WO 2001044276	A2	20010621	WO 2000-EP12698	20001214
WO 2001044276	A3	20011108		
W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR,				

HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG

BR 2000016432 A 20020917 BR 2000-16432 20001214
 EP 1244696 A2 20021002 EP 2000-983319 20001214

R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL, TR

US 2003157592 A1 20030821 US 2002-149759 20020613
 US 1999-171121P P 19991216
 WO 2000-EP12698 W 20001214

PRIORITY APPLN. INFO.:

AB Isolated nucleic acid mols., designated TCMRP (Tocopherol and ***Carotenoid*** Metab. Related Protein) nucleic acid mols., which encode novel TCMRPs from e.g. Physcomitrella patens are described. The invention also provides antisense nucleic acid mols., recombinant expression vectors contg. TCMRP nucleic acid mols., and host cells into which the expression vectors have been introduced. The invention still further provides isolated TCMRPs, mutated TCMRPs, fusion proteins, antigenic peptides and methods for the improvement of prodn. of a desired compd. from ***transformed*** cells, organisms or ***plants*** based on genetic engineering of TCMRP genes in these organisms.

L5 ANSWER 12 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 1999:127037 CAPLUS
 DOCUMENT NUMBER: 130:194482
 TITLE: Methods for producing transgenic ***plants*** and seeds with altered xanthophyll compositions
 INVENTOR(S): Shewmaker, Christine K.
 PATENT ASSIGNEE(S): Calgene LLC, USA
 SOURCE: PCT Int. Appl., 93 pp.
 CODEN: PIXXD2
 DOCUMENT TYPE: Patent
 LANGUAGE: English
 FAMILY ACC. NUM. COUNT: 2
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9907867	A1	19990218	WO 1998-US16466	19980806
W: AU, CA, CN, JP				
RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
US 6429356	B1	20020806	US 1997-908758	19970808
AU 9889002	A1	19990301	AU 1998-89002	19980806
AU 747542	B2	20020516		
EP 1002117	A1	20000524	EP 1998-940812	19980806
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, FI				
JP 2001512688	T2	20010828	JP 2000-506350	19980806
US 2002092039	A1	20020711	US 2002-41472	20020110
PRIORITY APPLN. INFO.:				
			US 1997-908758	A 19970808
			US 1996-24145P	P 19960809
			WO 1998-US16466	W 19980806
AB Methods are provided for producing ***plants*** and seeds having				

altered ***carotenoid*** compns. by ***transforming*** host
 plants with constructs having a transcriptional initiation region
 from a gene expressed in a ***plant*** seed, a plastid transit
 peptide, a DNA sequence derived from at least one ***carotenoid***
 biosynthesis gene coding region, and a transcriptional termination region.
 The methods find particular use in increasing the ***carotenoid***
 content in oilseed ***plants***. Transgenic Brassica napus, cotton,
 and Arabidopsis thaliana expressing the ***phytoene***

synthase gene crtB, ***phytoene*** ***desaturase*** gene
 crtI, geranylgeranyl pyrophosphate ***synthase*** gene crtE of Erwinia
 uredovora; or the .beta.-carotene hydroxylase gene crtZ or .beta.-carotene
 ketolase gene crtW of Agrobacterium aurantiacum; or both the crtB and crtI
 genes; or both the crtB and antisense lycopene .epsilon.-cyclase genes; or
 both antisense lycopene .beta.-cyclase and crtB genes; or both crtZ and
 crtB genes; or both crtW and crtB genes were prepd. These genes were
 fused to the napin transcription control region and the SSU leader
 sequence. The effects on ***carotenoid*** levels were detd.: all
 transgenic ***plants*** /seeds exhibited increased carotenoid levels.

REFERENCE COUNT: 9 THERE ARE 9 CITED REFERENCES AVAILABLE FOR THIS
 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L5 ANSWER 13 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 1998:126371 CAPLUS

DOCUMENT NUMBER: 128:203178

TITLE: Using enzymes of ***carotenoid*** biosynthesis to
 alter the ***carotenoid*** content and fatty acid
 profile of seeds

INVENTOR(S): Shewmaker, Christine K.

PATENT ASSIGNEE(S): Calgene, Inc., USA; Shewmaker, Christine K.

SOURCE: PCT Int. Appl., 70 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9806862	A1	19980219	WO 1997-US14035	19970808
W:	AL, AU, BA, BB, BG, BR, CA, CN, CU, CZ, EE, GE, HU, IL, IS, JP, KP, KR, LC, LK, LR, LT, LV, MG, MK, MN, MX, NO, NZ, PL, RO, SG, SI, SK, SL, TR, TT, UA, US, UZ, VN, YU, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
RW:	GH, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG			
AU 9740584	A1	19980306	AU 1997-40584	19970808
EP 925366	A1	19990630	EP 1997-938203	19970808
R:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, FI			
CN 1227609	A	19990901	CN 1997-197150	19970808
BR 9713462	A	20000328	BR 1997-13462	19970808
JP 2001505409	T2	20010424	JP 1998-509911	19970808
ZA 9707469	A	19980219	ZA 1997-7469	19970820
US 2002092039	A1	20020711	US 2002-41472	20020110
PRIORITY APPLN. INFO.:			US 1996-24145P P	19960809
			US 1997-908758 A1	19970808

AB Methods of altering the ***carotenoid*** content and fatty acid profile of seeds by altering the levels of expression of genes for enzymes of ***carotenoid*** biosynthesis. Increasing the diversion of acetate to ***carotenoid*** biosynthesis increases the anti-oxidant content of the oil, lowers the level of oxidn.-prone unsatd. fatty acids such as linoleate or linolenate, and increases the oleic acid content of the oil. Preferably, the enzyme is one of the earlier enzymes in the ***carotenoid*** pathway. The crtB gene of Erwinia uredovora, encoding ***phytoene*** ***synthase***, was placed under control of a napin gene promoter using the signal sequence of the RuBisCo small subunit gene and the construct introduced into Brassica napus by Agrobacterium-mediated ***transformation***. T2 ***plants*** showed Mendelian segregation of an orange phenotype. Seed from these ***plants*** showed increased levels of ***carotenoids*** and tocopherols, with several ***carotenoids*** not detectable in control seeds being found in transgenic seed. The fatty acid compn. of the seeds showed an increase in oleic acid content at the expense of linoleic and linolenic acid levels. Transgenic seeds showed slower germination than control seeds, but the germination rate was not affected.

REFERENCE COUNT: 16 THERE ARE 16 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L5 ANSWER 14 OF 14 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 1997:690574 CAPLUS

DOCUMENT NUMBER: 128:19146

TITLE: The regulation and genetic manipulation of
carotenoid biosynthesis in tomato fruit

AUTHOR(S): Bramley, Peter M.

CORPORATE SOURCE: Div. Biochem., Sch. Biol. Sci., Univ. London,
Egham/Surrey, TW20 0EX, UK

SOURCE: Pure and Applied Chemistry (1997), 69(10), 2159-2162
CODEN: PACHAS; ISSN: 0033-4545

PUBLISHER: Blackwell

DOCUMENT TYPE: Journal

LANGUAGE: English

AB The Ailsa Craig variety of tomato (Lycopersicon esculentum) has been ***transformed*** with ***carotenoid*** genes from higher ***plants*** and bacteria. Progeny have been analyzed for their ***carotenoid*** levels, carotenogenic enzyme activities and levels of gene expression. Ultrastructural studies have revealed changes in plastid structure. A similar approach has also been adopted with the high pigment (hp) mutant variety, which has elevated levels of ***carotenoids*** compared with the parental cultivar.

REFERENCE COUNT: 11 THERE ARE 11 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

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---Logging off of STN---

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Executing the logoff script...